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NEWS	1		Web Page URLs for STN Seminar Schedule - N. America
NEWS	2		"Ask CAS" for self-help around the clock
NEWS	3	SEP 09	CA/CAPLUS records now contain indexing from 1907 to the present
NEWS	4	Jul 15	Data from 1960-1976 added to RDISCLOSURE
NEWS	5	Jul 21	Identification of STN records implemented
NEWS	6	Jul 21	Polymer class term count added to REGISTRY
NEWS	7	Jul 22	INPADOC: Basic index (/BI) enhanced; Simultaneous Left and Right Truncation available
NEWS	8	AUG 05	New pricing for EUROPATFULL and PCTFULL effective August 1, 2003
NEWS	9	AUG 13	Field Availability (/FA) field enhanced in BEILSTEIN
NEWS	10	AUG 15	PATDPAFULL: one FREE connect hour, per account, in September 2003
NEWS	11	AUG 15	PCTGEN: one FREE connect hour, per account, in September 2003
NEWS	12	AUG 15	RDISCLOSURE: one FREE connect hour, per account, in September 2003
NEWS	13	AUG 15	TEMA: one FREE connect hour, per account, in September 2003
NEWS	14	AUG 18	Data available for download as a PDF in RDISCLOSURE
NEWS	15	AUG 18	Simultaneous left and right truncation added to PASCAL
NEWS	16	AUG 18	FROSTI and KOSMET enhanced with Simultaneous Left and Right Truncation
NEWS	17	AUG 18	Simultaneous left and right truncation added to ANABSTR
NEWS	18	SEP 22	DIPPR file reloaded
NEWS	19	SEP 25	INPADOC: Legal Status data to be reloaded
NEWS	20	SEP 29	DISSABS now available on STN
NEWS EXPRESS			April 4 CURRENT WINDOWS VERSION IS V6.01a, CURRENT MACINTOSH VERSION IS V6.0b(ENG) AND V6.0Jb(JP), AND CURRENT DISCOVER FILE IS DATED 01 APRIL 2003
NEWS HOURS			STN Operating Hours Plus Help Desk Availability
NEWS INTER			General Internet Information
NEWS LOGIN			Welcome Banner and News Items
NEWS PHONE			Direct Dial and Telecommunication Network Access to STN
NEWS WWW			CAS World Wide Web Site (general information)

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* * * * * STN Columbus * * * * *

FILE 'HOME' ENTERED AT 10:30:55 ON 30 SEP 2003

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COST IN U.S. DOLLARS

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FILE 'MEDLINE' ENTERED AT 10:31:25 ON 30 SEP 2003

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=> s hawkins phillip /au
L1 13 HAWKINS PHILLIP

=> s murry lynn /au
L2 1 MURRY LYNN

=> s GIPL
L3 218 GIPL

=> dup rem l1
PROCESSING COMPLETED FOR L1
L4 7 DUP REM L1 (6 DUPLICATES REMOVED)

=> d l4 total ibib

L4 ANSWER 1 OF 7 MEDLINE on STN DUPLICATE 1
ACCESSION NUMBER: 2002376138 MEDLINE
DOCUMENT NUMBER: 22117189 PubMed ID: 12121613
TITLE: Activation of phosphoinositide 3-kinase gamma by Ras.
AUTHOR: Suire Sabine; **Hawkins Phillip**; Stephens Len
CORPORATE SOURCE: Inositide Laboratory, The Babraham Institute, CB2 4AT
Cambridge, United Kingdom.
SOURCE: CURRENT BIOLOGY, (2002 Jul 9) 12 (13) 1068-75.
Journal code: 9107782. ISSN: 0960-9822.
PUB. COUNTRY: England: United Kingdom
DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
LANGUAGE: English
FILE SEGMENT: Priority Journals
ENTRY MONTH: 200302
ENTRY DATE: Entered STN: 20020718
Last Updated on STN: 20030205
Entered Medline: 20030204

L4 ANSWER 2 OF 7 MEDLINE on STN DUPLICATE 2
ACCESSION NUMBER: 2002159324 MEDLINE
DOCUMENT NUMBER: 21888450 PubMed ID: 11891120
TITLE: Roles of PI3Ks in leukocyte chemotaxis and phagocytosis.
AUTHOR: Stephens Len; Ellson Chris; **Hawkins Phillip**
CORPORATE SOURCE: The Babraham Institute, Babraham, Cambridge CB2 4AT, UK..
len.stephens@bbsrc.ac.uk
SOURCE: CURRENT OPINION IN CELL BIOLOGY, (2002 Apr) 14 (2) 203-13.
Ref: 53
Journal code: 8913428. ISSN: 0955-0674.

PUB. COUNTRY: United States
 DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
 General Review; (REVIEW)
 (REVIEW, ACADEMIC)
 LANGUAGE: English
 FILE SEGMENT: Priority Journals
 ENTRY MONTH: 200207
 ENTRY DATE: Entered STN: 20020314
 Last Updated on STN: 20020727
 Entered Medline: 20020726

L4 ANSWER 3 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN
 ACCESSION NUMBER: 1995:501942 CAPLUS
 DOCUMENT NUMBER: 122:231732
 TITLE: PDGF stimulates an increase in GTP-Rac via activation
 of phosphoinositide 3-kinase
 AUTHOR(S): **Hawkins, Phillip**; Eguinoa, Alicia; Qiu,
 Rong-Guo; Stokoe, David; Cooke, Frank T.; Walters,
 Rhodri; Wennstroem, Stefan; Claesson-Welsh, Lena;
 Evans, Tony; et al.
 CORPORATE SOURCE: The Babraham Inst., Cambridge, CB2 4AT, UK
 SOURCE: Current Biology (1995), 5(4), 393-403
 CODEN: CUBLE2; ISSN: 0960-9822
 PUBLISHER: Current Biology
 DOCUMENT TYPE: Journal
 LANGUAGE: English

L4 ANSWER 4 OF 7 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
 DUPLICATE 3
 ACCESSION NUMBER: 1995:154224 BIOSIS
 DOCUMENT NUMBER: PREV199598168524
 TITLE: Activation of the small GTP-binding proteins rho and rac by
 growth factor receptors.
 AUTHOR(S): Nobes, Catherine D.; **Hawkins, Phillip**; Stephens,
 Lens; Hall, Alan (1)
 CORPORATE SOURCE: (1) CRC Signal Transduction Oncogene Group, MRC Lab.
 Molecular Cell Biol., Univ. Coll. London, London WC1E 6BT
 UK
 SOURCE: Journal of Cell Science, (1995) Vol. 108, No. 1, pp.
 225-233.
 ISSN: 0021-9533.
 DOCUMENT TYPE: Article
 LANGUAGE: English

L4 ANSWER 5 OF 7 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. on STN
 DUPLICATE 4
 ACCESSION NUMBER: 1994:296238 BIOSIS
 DOCUMENT NUMBER: PREV199497309238
 TITLE: Activation of phosphoinositide 3-kinase is required for
 PDGF-stimulated membrane ruffling.
 AUTHOR(S): Wennstrom, Stefan; **Hawkins, Phillip**; Cooke,
 Frank; Hara, Kenta; Yonezawa, Kazuyoshi; Kasuga, Masato;
 Jackson, Trevor; Claesson-Welsh, Lena; Stephens, Len (1)
 CORPORATE SOURCE: (1) Dep. Dev. Signalling, AFRC Babraham Inst., Cambridge
 CB2 4AT UK
 SOURCE: Current Biology, (1994) Vol. 4, No. 5, pp. 385-393.
 ISSN: 0960-9822.
 DOCUMENT TYPE: Article
 LANGUAGE: English

L4 ANSWER 6 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN
 ACCESSION NUMBER: 1991:469063 CAPLUS
 DOCUMENT NUMBER: 115:69063
 TITLE: Inositol hexakisphosphate-membranes interactions: the

role of metal ions
AUTHOR(S): Cooke, Frank; Poyner, David; **Hawkins, Phillip**
; Erlebach, Christopher B.; Hanley, Michael
CORPORATE SOURCE: Lab. Mol. Biol., MRC Cent., Cambridge, CB2 2QH, UK
SOURCE: Biochemical Society Transactions (1991), 19(2), 152S
CODEN: BCSTB5; ISSN: 0300-5127
DOCUMENT TYPE: Journal
LANGUAGE: English

L4 ANSWER 7 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN
ACCESSION NUMBER: 1990:418445 CAPLUS
DOCUMENT NUMBER: 113:18445
TITLE: Phosphatidylinositol-3-phosphate and inositol
phosphates in mitogenesis
AUTHOR(S): Poyner, David; **Hawkins, Phillip**; Hanley,
Michael
CORPORATE SOURCE: Med. Sch., Univ. Cambridge, Cambridge, CB2 2QH, UK
SOURCE: Biochemical Society Transactions (1990), 18(3), 450-1
CODEN: BCSTB5; ISSN: 0300-5127
DOCUMENT TYPE: Journal
LANGUAGE: English

Sequence Comparison A

RESULT 10
PCT-US93-00643-11
; Sequence 11, Application PC/TUS9300643
; GENERAL INFORMATION:
; APPLICANT: Baylink, David J.
; APPLICANT: Linkhart, Susan
; TITLE OF INVENTION: AMINO PROCOLLAGEN 1(I) PEPTIDES
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend
; STREET: One Market Plaza, Steuart Street Tower
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94105-1492
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US93/00643
; FILING DATE: 19930125
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parmelee, Steven W.
; REGISTRATION NUMBER: 31,990
; REFERENCE/DOCKET NUMBER: 14508-3
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 467-9600
; TELEFAX: (415) 543-5043
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US93-00643-11

Query Match 2.9%; Score 6; DB 5; Length 160;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 182 PGA EVP 187
|||
Db 82 PGA EVP 87

09875520 Results

SEQ ID NO: 2 oligo

SUMMARIES

Result No.	Score	% Match	Query Length	DB	ID	Description
1	204	100.0	204	18	AAW26579	Novel phospholipas
2	204	100.0	204	19	AAW79142	Human phospholipas
3	204	100.0	204	20	AAV30147	Amino acid sequenc
4	73	35.8	89	21	AAG03252	Human secreted pro
5	8	3.9	73	22	AAO12980	Human polypeptide
6	8	3.9	125	22	AAG76150	Human colon cancer
7	8	3.9	304	22	ABG19393	Novel human diagno
8	8	3.9	314	22	ABG10498	Novel human diagno
9	8	3.9	315	21	AAB58262	Lung cancer associ

Result No.	Score	% Match	Query Length	DB	ID	Description
1	8	3.9	133	2	B82977	conserved hypothet
2	8	3.9	340	2	T28686	hypothetical prote
3	8	3.9	436	2	T28066	hypothetical prote
4	7	3.4	114	2	JC5860	polyketide synthas
5	7	3.4	144	2	D83864	hypothetical prote
6	7	3.4	150	2	A69551	conserved hypothet
7	7	3.4	151	2	T34245	hypothetical prote
8	7	3.4	185	2	S12676	hypothetical prote
9	7	3.4	193	2	C75374	hypothetical prote
10	7	3.4	236	2	A84823	xanthine phosphori
11	7	3.4	254	2	T30675	hypothetical prote
12	7	3.4	258	2	S69056	probable 28k struc
13	7	3.4	309	2	F83605	histone H1 - yeast
14	7	3.4	316	2	F87260	probable permease
15	7	3.4	323	2	E95864	WecB/TagA/CpsF fam
16	7	3.4	331	2	T18247	probable ABC trans
17	7	3.4	336	2	E96814	transcription regu
18	7	3.4	360	2	S36750	hypothetical prote
19	7	3.4	422	2	T12786	cannabinoid recept
20	7	3.4	520	2	S74497	conserved hypothet
						hypothetical prote

SUMMARIES

Result No.	Score	% Match	Query Length	DB	ID	Description
1	7	3.4	185	1	YALI_TRYBB	P17960 trypanosoma
2	7	3.4	258	1	H1_YEAST	P53551 saccharomyc
3	7	3.4	360	1	CB2R_HUMAN	P34972 homo sapien
4	7	3.4	814	1	SLA1_BACAN	P49051 bacillus an
5	7	3.4	2560	1	PPS2_BACSU	P39846 bacillus su
6	7	3.4	2869	1	RBP1_PLAVB	Q00798 plasmodium
7	6	2.9	35	1	SCKK_TITSE	P56219 tityus serr
9	6	2.9	54	1	PSBK_EUGGR	P31481 euglena gra
10	6	2.9	54	1	PSBK_EUGST	Q9ms58 euglena ste
11	6	2.9	55	1	PSBK_MARPO	P10348 marchantia
12	6	2.9	56	1	PSBK_PINTH	P41598 pinus thunb
13	6	2.9	58	1	RL30_ECOLI	P02430 escherichia
14	6	2.9	58	1	RL30_SALTY	O54300 salmonella
15	6	2.9	59	1	PSBK_SPIOL	P12163 spinacia ol
16	6	2.9	59	1	RL30_BUCAK	P46184 buchnera ap
17	6	2.9	61	1	PSBK_HORVU	P25877 hordeum vul
18	6	2.9	61	1	PSBK_LOTJA	Q9bbs2 lotus

SUMMARIES

Result	Query						
No.	Score	Match	Length	DB	ID		Description
1	141	69.1	254	4	O95053		O95053 homo sapien
2	12	5.9	212	11	Q9D6V5		Q9d6v5 mus musculu
3	12	5.9	212	11	Q9CQD7		Q9cq d7 mus musculu
4	8	3.9	133	16	Q9HTK6		Q9htk6 pseudomonas
5	8	3.9	340	16	O69810		O69810 streptomyce
6	8	3.9	436	5	Q23653		Q23653 caenorhabdi
7	8	3.9	801	16	Q9A0I0		Q9a0i0 streptococc
8	7	3.4	114	2	O32461		O32461 actinomadur
9	7	3.4	118	6	Q95JV4		Q95jv4 macaca fasc
10	7	3.4	133	12	Q83891		Q83891 ovine adeno
11	7	3.4	144	16	Q9KC58		Q9kc58 bacillus ha
12	7	3.4	150	17	O30263		O30263 archaeoglob
13	7	3.4	151	5	Q19809		Q19809 caenorhabdi
14	7	3.4	184	5	P90604		P90604 trypanosoma
15	7	3.4	193	2	Q47381		Q47381 escherichia
16	7	3.4	193	16	Q9RTY2		Q9rty2 deinococcus
17	7	3.4	219	16	Q92Q39		Q92q39 rhizobium m
18	7	3.4	254	12	Q98241		Q98241 molluscum c
19	7	3.4	283	10	O04195		O04195 arabidopsis

RESULT 7

US-08-318-193-16

; Sequence 16, Application US/08318193

; Patent No. 5641663

; GENERAL INFORMATION:

; APPLICANT: GARVIN, Robert T.

; APPLICANT: MALEK, Lawrence T.

; TITLE OF INVENTION: AN EXPRESSION SYSTEM FOR THE SECRETION

; TITLE OF INVENTION: OF BIOACTIVE HUMAN GRANULOCYTE MACROPHAGE COLONY

; TITLE OF INVENTION: STIMULATING FACTOR (GM-CSF) AND OTHER HETEROLOGOUS

; TITLE OF INVENTION: PROTEINS FROM STREPTOMYCES

; NUMBER OF SEQUENCES: 91

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Foley & Lardner

; STREET: 1800 Diagonal Road, Suite 500

; CITY: Alexandria

; STATE: Virginia

; COUNTRY: USA

; ZIP: 22313-0299

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/318,193

; FILING DATE:

; CLASSIFICATION: 435

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/07/935,314

; FILING DATE:

; APPLICATION NUMBER: US 07/224,568

; ATTORNEY/AGENT INFORMATION:

; NAME: BENT, Stephen A.

; REGISTRATION NUMBER: 29,768

; REFERENCE/DOCKET NUMBER: 18740/116 CACO

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: (703)836-9300

; TELEFAX: (703)683-4109

; TELEX: 899149

; INFORMATION FOR SEQ ID NO: 16:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 144 amino acids

; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-318-193-16

Query Match 2.9%; Score 6; DB 1; Length 144;
Best Local Similarity 100.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 113 SVPLTN 118
|||||
Db 85 SVPLTN 90

RESULT 10
PCT-US93-00643-11
; Sequence 11, Application PC/TUS9300643
; GENERAL INFORMATION:
; APPLICANT: Baylink, David J.
; APPLICANT: Linkhart, Susan
; TITLE OF INVENTION: AMINO PROCOLLAGEN 1(I) PEPTIDES
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend
; STREET: One Market Plaza, Steuart Street Tower
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94105-1492
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US93/00643
; FILING DATE: 19930125
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parmelee, Steven W.
; REGISTRATION NUMBER: 31,990
; REFERENCE/DOCKET NUMBER: 14508-3
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 467-9600
; TELEFAX: (415) 543-5043
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US93-00643-11

Query Match 2.9%; Score 6; DB 5; Length 160;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 182 PGA EVP 187
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Db 82 PGA EVP 87

RESULT 9
US-08-479-233-11
; Sequence 11, Application US/08479233
; Patent No. 5599679
; GENERAL INFORMATION:
; APPLICANT: Baylink, David J.


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; APPLICANT: Linkhart, Susan
; TITLE OF INVENTION: AMINO PROCOLLAGEN 1(I) PEPTIDE
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend
; STREET: One Market Plaza, Steuart Street Tower
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94105-1492
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/479,233
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/07/829,142
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Parmelee, Steven W.
; REGISTRATION NUMBER: 31,990
; REFERENCE/DOCKET NUMBER: 14508-3
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 467-9600
; TELEFAX: (415) 543-5043
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-479-233-11

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Query Match          2.9%; Score 6; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy 182 PGA EVP 187
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Db  82 PGA EVP 87

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SEQ ID NO: 2

SUMMARIES

Result No.	Score	% Query				Description
		Match	Length	DB	ID	
1	1128	100.0	204	18	AAW26579	Novel phospholipas
2	1128	100.0	204	19	AAW79142	Human phospholipas
3	1128	100.0	204	20	AAY30147	Amino acid sequenc
4	493	43.7	89	21	AAG03252	Human secreted pro
5	224	19.9	181	20	AAY26145	Phospholipase A2 i
6	223	19.8	181	20	AAY26138	Phospholipase A2 i
7	223	19.8	181	20	AAY26113	Phospholipase A2 i
8	223	19.8	181	21	AAY83613	Mature beta chain

9	222	19.7	181	20	AAY26151	Phospholipase A2 i
10	212	18.8	181	20	AAY26159	Phospholipase A2 i
11	211	18.7	181	20	AAY26158	Phospholipase A2 i
12	204.5	18.1	182	20	AAY26152	Phospholipase A2 i
13	186.5	16.5	182	20	AAY26165	Phospholipase A2 i
14	167	14.8	220	23	AAU10477	Mouse pancreatic p
15	162	14.4	201	22	AAE03680	Python reticulatus
16	156	13.8	182	22	AAE03682	Python reticulatus
17	136	12.1	335	21	AAB26241	Human urokinase-ty
18	135	12.0	299	22	AAU07611	Human u-PAR deleti
19	135	12.0	335	7	AAP60436	Sequence of human
20	135	12.0	335	11	AAR07561	Recombinant urokin
21	135	12.0	335	14	AAR44424	Human phospholipas
22	135	12.0	335	15	AAR58707	Human phospholipas
23	135	12.0	335	17	AAR97612	Human urokinase pl
24	135	12.0	335	18	AAW31165	Human phospholipas
25	135	12.0	335	20	AAY04103	Urokinase-type pla
26	135	12.0	335	22	AAU04454	Human urokinase-ty
27	135	12.0	335	22	AAU07610	Human u-PAR substi
28	132.5	11.7	202	21	AAY83648	NSI Phospholipase
29	132	11.7	202	21	AAY83582	Phospholipase A_2
30	128.5	11.4	183	20	AAY26143	Phospholipase A2 i
31	128	11.3	231	22	AAM25880	Human protein sequ
32	126	11.2	182	20	AAY26111	Phospholipase A2 i
33	125.5	11.1	183	20	AAY26135	Phospholipase A2 i
34	125.5	11.1	183	20	AAY26157	Phospholipase A2 i
35	125.5	11.1	183	21	AAY83610	Phospholipase A2 i
36	125	11.1	202	21	AAY83583	Mature alpha chain
37	124.5	11.0	183	20	AAY26137	Phospholipase A_2
38	124.5	11.0	183	20	AAY26144	Phospholipase A2 i
39	124.5	11.0	183	20	AAY26149	Phospholipase A2 i
40	124.5	11.0	183	20	AAY26150	Phospholipase A2 i
41	124.5	11.0	183	20	AAY26112	Phospholipase A2 i
42	124.5	11.0	183	21	AAY83612	Phospholipase A2 i
43	124	11.0	182	20	AAY26110	Mature alpha chain
44	121.5	10.8	237	20	AAY02654	Phospholipase A2 i
45	120.5	10.7	182	20	AAY26136	Human secreted pro
						Phospholipase A2 i

Issued:

Result No.	Score	Query Match	Length	DB	ID	Description
1	1128	100.0	204	1	US-08-652-859-2	Sequence 2, Appli
2	1128	100.0	204	2	US-08-919-706-2	Sequence 2, Appli
3	1128	100.0	204	2	US-09-153-751-2	Sequence 2, Appli
4	171	15.2	200	1	US-08-652-859-3	Sequence 3, Appli
5	171	15.2	200	2	US-08-919-706-3	Sequence 3, Appli
6	171	15.2	200	2	US-09-153-751-3	Sequence 3, Appli
7	135	12.0	335	1	US-08-085-122-13	Sequence 13, Appl
8	135	12.0	335	4	US-08-442-108B-23	Sequence 23, Appl
9	121.5	10.8	237	4	US-09-227-357-154	Sequence 154, App
10	114	10.1	437	3	US-09-073-569-2	Sequence 2, Appli
11	99	8.8	123	4	US-09-203-939-4	Sequence 4, Appli
12	99	8.8	123	4	US-09-251-835-4	Sequence 4, Appli
13	99	8.8	123	4	US-09-318-503-4	Sequence 4, Appli
14	99	8.8	123	4	US-09-038-261A-4	Sequence 4, Appli
15	97	8.6	123	4	US-09-203-939-7	Sequence 7, Appli
16	97	8.6	123	4	US-09-251-835-7	Sequence 7, Appli
17	97	8.6	123	4	US-09-318-503-7	Sequence 7, Appli
18	97	8.6	123	4	US-09-038-261A-7	Sequence 7, Appli
19	96.5	8.6	1917	4	US-09-627-650B-5	Sequence 5, Appli
20	96.5	8.6	1917	4	US-09-436-063C-5	Sequence 5,

Result No.	Score	Query Match Length	DB	ID	Description
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1	228	20.2	185	1	JC2394	phospholipase A2 i
2	171	15.2	200	1	A54020	Crotalus neutraliz
3	165	14.6	330	1	JN0561	urokinase-type pla
4	141	12.5	188	1	JC2393	phospholipase A2 i
5	135	12.0	335	2	A39743	u-plasminogen acti
6	119.5	10.6	327	2	A55356	urokinase-type pla
7	118	10.5	328	2	S42152	urinary plasminoge
8	118	10.5	5376	2	T42215	zonadhesin - mouse
9	117	10.4	126	2	S53340	CD59 protein - rat
10	113	10.0	1101	2	T16840	hypothetical prote
11	111	9.8	506	2	A40679	transcription enha
12	111	9.8	523	2	B40679	transcription enha
13	102.5	9.1	1360	2	T12064	DNA binding protei
14	102	9.0	2907	2	A57278	fibrillin-2 precur